

John S K Kauwe

List of Publications by Year in descending order

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Version: 2024-02-01

136
papers

24,787
citations

38660

50
h-index

11581

135
g-index

156
all docs

156
docs citations

156
times ranked

23824
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 1 | Analysis of high-risk pedigrees identifies 11 candidate variants for Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2022, 18, 307-317. | 0.4 | 10 |
| 2 | Association between WWOX/MAF variants and dementia-related neuropathologic endophenotypes. <i>Neurobiology of Aging</i> , 2022, 111, 95-106. | 1.5 | 6 |
| 3 | Knowledge Gaps, Challenges, and Opportunities in Health and Prevention Research for Asian Americans, Native Hawaiians, and Pacific Islanders: A Report From the 2021 National Institutes of Health Workshop. <i>Annals of Internal Medicine</i> , 2022, 175, 574-589. | 2.0 | 40 |
| 4 | Genome-wide association study of brain arteriolosclerosis. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2022, 42, 1437-1450. | 2.4 | 2 |
| 5 | New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436. | 9.4 | 700 |
| 6 | The Ramp Atlas: facilitating tissue and cell-specific ramp sequence analyses through an intuitive web interface. <i>NAR Genomics and Bioinformatics</i> , 2022, 4, . | 1.5 | 3 |
| 7 | Novel Alzheimer Disease Risk Loci and Pathways in African American Individuals Using the African Genome Resources Panel. <i>JAMA Neurology</i> , 2021, 78, 102. | 4.5 | 144 |
| 8 | GenoRisk: A polygenic risk score for Alzheimer's disease. <i>Alzheimer's and Dementia: Translational Research and Clinical Interventions</i> , 2021, 7, e12211. | 1.8 | 7 |
| 9 | A comprehensive analysis of the phylogenetic signal in ramp sequences in 211 vertebrates. <i>Scientific Reports</i> , 2021, 11, 622. | 1.6 | 2 |
| 10 | Alzheimer's disease alters oligodendrocytic glycolytic and ketolytic gene expression. <i>Alzheimer's and Dementia</i> , 2021, 17, 1474-1486. | 0.4 | 37 |
| 11 | Phylogeography of two marine predators, giant trevally (<i>Caranx ignobilis</i>) and bluefin trevally (<i>Caranx melampygus</i>), across the Indo-Pacific. <i>Bulletin of Marine Science</i> , 2021, 97, 257-280. | 0.4 | 6 |
| 12 | Alzheimer's Disease Alters Oligodendrocytic Glycolytic and Ketolytic Gene Expression. <i>FASEB Journal</i> , 2021, 35, . | 0.2 | 1 |
| 13 | <i>De novo</i> genome assembly of the marine teleost, bluefin trevally (<i>Caranx melampygus</i>). <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, . | 0.8 | 2 |
| 14 | Analysis of genes (TMEM106B, GRN, ABCC9, KCNMB2, and APOE) implicated in risk for LATE-NC and hippocampal sclerosis provides pathogenetic insights: a retrospective genetic association study. <i>Acta Neuropathologica Communications</i> , 2021, 9, 152. | 2.4 | 26 |
| 15 | Pairwise Correlation Analysis of the Alzheimer's Disease Neuroimaging Initiative (ADNI) Dataset Reveals Significant Feature Correlation. <i>Genes</i> , 2021, 12, 1661. | 1.0 | 5 |
| 16 | Interaction Between Physical Activity and Genes Related to Neurotrophin Signaling in Late-Life Cognitive Performance: The Cache County Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2020, 75, 1633-1642. | 1.7 | 7 |
| 17 | CUBAP: an interactive web portal for analyzing codon usage biases across populations. <i>Nucleic Acids Research</i> , 2020, 48, 11030-11039. | 6.5 | 7 |
| 18 | Lingering Taxonomic Challenges Hinder Conservation and Management of Global Bonefishes. <i>Fisheries</i> , 2020, 45, 347-358. | 0.6 | 15 |

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|----|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 19 | Distinct clinicopathologic clusters of persons with TDP-43 proteinopathy. <i>Acta Neuropathologica</i> , 2020, 140, 659-674. | 3.9 | 29 |
| 20 | Atypical chemokine receptor ACKR2-V41A has decreased CCL2 binding, scavenging, and activation, supporting sustained inflammation and increased Alzheimer's disease risk. <i>Scientific Reports</i> , 2020, 10, 8019. | 1.6 | 7 |
| 21 | Codon Pairs are Phylogenetically Conserved: A comprehensive analysis of codon pairing conservation across the Tree of Life. <i>PLoS ONE</i> , 2020, 15, e0232260. | 1.1 | 8 |
| 22 | Genome-wide association study of rate of cognitive decline in Alzheimer's disease patients identifies novel genes and pathways. <i>Alzheimer's and Dementia</i> , 2020, 16, 1134-1145. | 0.4 | 28 |
| 23 | Identification and genomic analysis of pedigrees with exceptional longevity identifies candidate rare variants. <i>Neurobiology of Disease</i> , 2020, 143, 104972. | 2.1 | 7 |
| 24 | Predicting Clinical Dementia Rating Using Blood RNA Levels. <i>Genes</i> , 2020, 11, 706. | 1.0 | 10 |
| 25 | Failure to detect synergy between variants in transferrin and hemochromatosis and Alzheimer's disease in large cohort. <i>Neurobiology of Aging</i> , 2020, 89, 142.e9-142.e12. | 1.5 | 9 |
| 26 | RAB10: an Alzheimer's disease resilience locus and potential drug target. <i>Clinical Interventions in Aging</i> , 2019, Volume 14, 73-79. | 1.3 | 37 |
| 27 | Systematic analysis of dark and camouflaged genes reveals disease-relevant genes hiding in plain sight. <i>Genome Biology</i> , 2019, 20, 97. | 3.8 | 122 |
| 28 | Association study of rs3846662 with Alzheimer's disease in a population-based cohort: the Cache County Study. <i>Neurobiology of Aging</i> , 2019, 84, 242.e1-242.e6. | 1.5 | 5 |
| 29 | Relative risk for Alzheimer disease based on complete family history. <i>Neurology</i> , 2019, 92, e1745-e1753. | 1.5 | 45 |
| 30 | Association of Rare Coding Mutations With Alzheimer Disease and Other Dementias Among Adults of European Ancestry. <i>JAMA Network Open</i> , 2019, 2, e191350. | 2.8 | 58 |
| 31 | Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β , tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019, 51, 414-430. | 9.4 | 1,962 |
| 32 | Assembly of 809 whole mitochondrial genomes with clinical, imaging, and fluid biomarker phenotyping. <i>Alzheimer's and Dementia</i> , 2018, 14, 514-519. | 0.4 | 14 |
| 33 | Mitochondria and Alzheimer's Disease: the Role of Mitochondrial Genetic Variation. <i>Current Genetic Medicine Reports</i> , 2018, 6, 1-10. | 1.9 | 45 |
| 34 | Common DNA Variants Accurately Rank an Individual of Extreme Height. <i>International Journal of Genomics</i> , 2018, 2018, 1-7. | 0.8 | 5 |
| 35 | Genome-wide association study for variants that modulate relationships between cerebrospinal fluid amyloid-beta 42, tau, and p-tau levels. <i>Alzheimer's Research and Therapy</i> , 2018, 10, 86. | 3.0 | 18 |
| 36 | Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, . | 6.0 | 1,085 |

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|----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 37 | Population genealogy resource shows evidence of familial clustering for Alzheimer disease. <i>Neurology: Genetics</i> , 2018, 4, e249. | 0.9 | 6 |
| 38 | A Common Variant of IL-6R is Associated with Elevated IL-6 Pathway Activity in Alzheimer's Disease Brains. <i>Journal of Alzheimer's Disease</i> , 2017, 56, 1037-1054. | 1.2 | 44 |
| 39 | Systems biology approach to late-onset Alzheimer's disease genome-wide association study identifies novel candidate genes validated using brain expression data and <i>Caenorhabditis elegans</i> experiments. , 2017, 13, 1133-1142. | | 40 |
| 40 | Genome-wide association study identifies four novel loci associated with Alzheimer's endophenotypes and disease modifiers. <i>Acta Neuropathologica</i> , 2017, 133, 839-856. | 3.9 | 199 |
| 41 | Transethnic genome-wide scan identifies novel Alzheimer's disease loci. <i>Alzheimer's and Dementia</i> , 2017, 13, 727-738. | 0.4 | 166 |
| 42 | CSF protein changes associated with hippocampal sclerosis risk gene variants highlight impact of GRN/PGRN. <i>Experimental Gerontology</i> , 2017, 90, 83-89. | 1.2 | 7 |
| 43 | Sex Differences in Risk for Alzheimer's Disease Related to Neurotrophin Gene Polymorphisms: The Cache County Memory Study. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2017, 72, 1607-1613. | 1.7 | 15 |
| 44 | A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. <i>Nature Neuroscience</i> , 2017, 20, 1052-1061. | 7.1 | 330 |
| 45 | Using the Health Belief Model to evaluate Samoan caregiver perceptions for rheumatic heart disease follow-up care. <i>International Journal of Health Promotion and Education</i> , 2017, 55, 148-157. | 0.4 | 2 |
| 46 | Genome-wide, high-content siRNA screening identifies the Alzheimer's genetic risk factor FERMT2 as a major modulator of APP metabolism. <i>Acta Neuropathologica</i> , 2017, 133, 955-966. | 3.9 | 60 |
| 47 | Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384. | 9.4 | 783 |
| 48 | [O11]: CEREBROSPINAL FLUID ENDOPHENOTYPES PROVIDE INSIGHT INTO BIOLOGY UNDERLYING ALZHEIMER'S DISEASE. <i>Alzheimer's and Dementia</i> , 2017, 13, P218. | 0.4 | 0 |
| 49 | Linkage, whole genome sequence, and biological data implicate variants in RAB10 in Alzheimer's disease resilience. <i>Genome Medicine</i> , 2017, 9, 100. | 3.6 | 67 |
| 50 | The Opioid Abuse Risk Screener predicts aberrant same-day urine drug tests and 1-year controlled substance database checks: A brief report. <i>Health Psychology Open</i> , 2017, 4, 205510291774845. | 0.7 | 2 |
| 51 | Fine-mapping of the human leukocyte antigen locus as a risk factor for Alzheimer disease: A case-control study. <i>PLoS Medicine</i> , 2017, 14, e1002272. | 3.9 | 67 |
| 52 | Discovery and Confirmation of Diagnostic Serum Lipid Biomarkers for Alzheimer's Disease Using Direct Infusion Mass Spectrometry. <i>Journal of Alzheimer's Disease</i> , 2017, 59, 277-290. | 1.2 | 19 |
| 53 | Seroprevalence and Serointensity of Latent <i>Toxoplasma gondii</i> in a Sample of Elderly Adults With and Without Alzheimer Disease. <i>Alzheimer Disease and Associated Disorders</i> , 2016, 30, 123-126. | 0.6 | 33 |
| 54 | Genetic studies of plasma analytes identify novel potential biomarkers for several complex traits. <i>Scientific Reports</i> , 2016, 6, . | 1.6 | 25 |

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|----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 55 | Genome-wide association study of prolactin levels in blood plasma and cerebrospinal fluid. <i>BMC Genomics</i> , 2016, 17, 436. | 1.2 | 2 |
| 56 | Assessment of the genetic variance of late-onset Alzheimer's disease. <i>Neurobiology of Aging</i> , 2016, 41, 200.e13-200.e20. | 1.5 | 174 |
| 57 | Crowdsourced estimation of cognitive decline and resilience in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2016, 12, 645-653. | 0.4 | 72 |
| 58 | Shared genetic contribution to ischemic stroke and Alzheimer's disease. <i>Annals of Neurology</i> , 2016, 79, 739-747. | 2.8 | 56 |
| 59 | Presenilin E318G variant and Alzheimer's disease risk: the Cache County study. <i>BMC Genomics</i> , 2016, 17, 438. | 1.2 | 11 |
| 60 | Evaluating the necessity of PCR duplicate removal from next-generation sequencing data and a comparison of approaches. <i>BMC Bioinformatics</i> , 2016, 17, 239. | 1.2 | 124 |
| 61 | Variants in ACPA are associated with cerebrospinal fluid Prostatic Acid Phosphatase levels. <i>BMC Genomics</i> , 2016, 17, 439. | 1.2 | 1 |
| 62 | Variants in CCL16 are associated with blood plasma and cerebrospinal fluid CCL16 protein levels. <i>BMC Genomics</i> , 2016, 17, 437. | 1.2 | 1 |
| 63 | Genes for a "Welllderly" Life. <i>Trends in Molecular Medicine</i> , 2016, 22, 637-639. | 3.5 | 7 |
| 64 | A potential endophenotype for Alzheimer's disease: cerebrospinal fluid clusterin. <i>Neurobiology of Aging</i> , 2016, 37, 208.e1-208.e9. | 1.5 | 44 |
| 65 | Interaction between variants in <i>CLU</i> and <i>MS4A4E</i> modulates Alzheimer's disease risk. <i>Alzheimer's and Dementia</i> , 2016, 12, 121-129. | 0.4 | 18 |
| 66 | A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , 2016, 21, 108-117. | 4.1 | 260 |
| 67 | Influence of Coding Variability in APP- $\text{A}\beta$ Metabolism Genes in Sporadic Alzheimer's Disease. <i>PLoS ONE</i> , 2016, 11, e0150079. | 1.1 | 34 |
| 68 | Discovery and Subsequent Confirmation of Novel Serum Biomarkers Diagnosing Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2015, 49, 317-327. | 1.2 | 10 |
| 69 | Bridging the Gap between Statistical and Biological Epistasis in Alzheimer's Disease. <i>BioMed Research International</i> , 2015, 2015, 1-7. | 0.9 | 26 |
| 70 | Rarity of the Alzheimer Disease "Protective" <i>APP</i> A673T Variant in the United States. <i>JAMA Neurology</i> , 2015, 72, 209. | 4.5 | 41 |
| 71 | Genetic studies of quantitative MCI and AD phenotypes in ADNI: Progress, opportunities, and plans. <i>Alzheimer's and Dementia</i> , 2015, 11, 792-814. | 0.4 | 241 |
| 72 | Genetically predicted body mass index and Alzheimer's disease-related phenotypes in three large samples: Mendelian randomization analyses. <i>Alzheimer's and Dementia</i> , 2015, 11, 1439-1451. | 0.4 | 46 |

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|----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 73 | Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 658-671. | 0.4 | 173 |
| 74 | Population-based analysis of cholesteryl ester transfer protein identifies association between I405V and cognitive decline: the Cache County Study. <i>Neurobiology of Aging</i> , 2015, 36, 547.e1-547.e3. | 1.5 | 8 |
| 75 | Genome-Wide Association Study of CSF Levels of 59 Alzheimer's Disease Candidate Proteins: Significant Associations with Proteins Involved in Amyloid Processing and Inflammation. <i>PLoS Genetics</i> , 2014, 10, e1004758. | 1.5 | 109 |
| 76 | A Versatile Omnibus Test for Detecting Mean and Variance Heterogeneity. <i>Genetic Epidemiology</i> , 2014, 38, 51-59. | 0.6 | 52 |
| 77 | Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. <i>JAMA Neurology</i> , 2014, 71, 1394. | 4.5 | 166 |
| 78 | Population-based Analysis of Alzheimer's Disease Risk Alleles Implicates Genetic Interactions. <i>Biological Psychiatry</i> , 2014, 75, 732-737. | 0.7 | 52 |
| 79 | Observed Changes in Radiographic Measurements of the First Ray after Frontal Plane Rotation of the First Metatarsal in a Cadaveric Foot Model. <i>Journal of Foot and Ankle Surgery</i> , 2014, 53, 274-278. | 0.5 | 40 |
| 80 | Genetic Discoveries in AD Using CSF Amyloid and Tau. <i>Current Genetic Medicine Reports</i> , 2014, 2, 23-29. | 1.9 | 10 |
| 81 | Genetic analysis of quantitative phenotypes in AD and MCI: imaging, cognition and biomarkers. <i>Brain Imaging and Behavior</i> , 2014, 8, 183-207. | 1.1 | 161 |
| 82 | Rare coding variants in the phospholipase D3 gene confer risk for Alzheimer's disease. <i>Nature</i> , 2014, 505, 550-554. | 13.7 | 425 |
| 83 | Variant Tool Chest: an improved tool to analyze and manipulate variant call format (VCF) files. <i>BMC Bioinformatics</i> , 2014, 15, S12. | 1.2 | 11 |
| 84 | Mitochondrial genomic variation associated with higher mitochondrial copy number: the Cache County Study on Memory Health and Aging. <i>BMC Bioinformatics</i> , 2014, 15, S6. | 1.2 | 15 |
| 85 | Population substructure in Cache County, Utah: the Cache County study. <i>BMC Bioinformatics</i> , 2014, 15, S8. | 1.2 | 7 |
| 86 | Observed Changes in First Metatarsal and Medial Cuneiform Positions after First Metatarsophalangeal Joint Arthrodesis. <i>Journal of Foot and Ankle Surgery</i> , 2014, 53, 32-35. | 0.5 | 12 |
| 87 | Phosphorylated Tau-A ²⁴² Ratio as a Continuous Trait for Biomarker Discovery for Early-Stage Alzheimer's Disease in Multiplex Immunoassay Panels of Cerebrospinal Fluid. <i>Biological Psychiatry</i> , 2014, 75, 723-731. | 0.7 | 72 |
| 88 | Missense variant in TREML2 protects against Alzheimer's disease. <i>Neurobiology of Aging</i> , 2014, 35, 1510.e19-1510.e26. | 1.5 | 110 |
| 89 | Variants in <i>PPP3R1</i> and <i>MAPT</i> are associated with more rapid functional decline in Alzheimer's disease: The Cache County Dementia Progression Study. <i>Alzheimer's and Dementia</i> , 2014, 10, 366-371. | 0.4 | 36 |
| 90 | Calibrating Longitudinal Cognition in Alzheimer's Disease Across Diverse Test Batteries and Datasets. <i>Neuroepidemiology</i> , 2014, 43, 194-205. | 1.1 | 43 |

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|-----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 91 | Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661. | 1.1 | 155 |
| 92 | Assessment of TREM2 rs75932628 association with Alzheimer's disease in a population-based sample: the Cache County Study. Neurobiology of Aging, 2013, 34, 2889.e11-2889.e13. | 1.5 | 47 |
| 93 | Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. Nature Genetics, 2013, 45, 1452-1458. | 9.4 | 3,741 |
| 94 | TREM2 Variants in Alzheimer's Disease. New England Journal of Medicine, 2013, 368, 117-127. | 13.9 | 2,385 |
| 95 | O4-01-01: Association of genetic variants with cerebrospinal fluid protein levels of ACE, MMP3 and other proteins and risk for Alzheimer's disease. , 2013, 9, P677-P678. | | 1 |
| 96 | GWAS of Cerebrospinal Fluid Tau Levels Identifies Risk Variants for Alzheimer's Disease. Neuron, 2013, 78, 256-268. | 3.8 | 344 |
| 97 | Genetics of Alzheimer's Disease. BioMed Research International, 2013, 2013, 1-13. | 0.9 | 75 |
| 98 | Mitochondrial Haplotypes Associated with Biomarkers for Alzheimer's Disease. PLoS ONE, 2013, 8, e74158. | 1.1 | 28 |
| 99 | Alzheimer's Disease: Analyzing the Missing Heritability. PLoS ONE, 2013, 8, e79771. | 1.1 | 257 |
| 100 | Strong Evidence for a Genetic Contribution to Late-Onset Alzheimer's Disease Mortality: A Population-Based Study. PLoS ONE, 2013, 8, e77087. | 1.1 | 14 |
| 101 | The Role of Variation at APP, PSEN1, PSEN2, and MAPT in Late Onset Alzheimer's Disease. Journal of Alzheimer's Disease, 2012, 28, 377-387. | 1.2 | 53 |
| 102 | Cerebrospinal fluid APOE levels: an endophenotype for genetic studies for Alzheimer's disease. Human Molecular Genetics, 2012, 21, 4558-4571. | 1.4 | 196 |
| 103 | Novel late-onset Alzheimer disease loci variants associate with brain gene expression. Neurology, 2012, 79, 221-228. | 1.5 | 144 |
| 104 | Single nucleotide polymorphism discovery in cutthroat trout subspecies using genome reduction, barcoding, and 454 pyro-sequencing. BMC Genomics, 2012, 13, 724. | 1.2 | 18 |
| 105 | Mitochondrial Genomic Analysis of Late Onset Alzheimer's Disease Reveals Protective Haplogroups H6A1A/H6A1B: The Cache County Study on Memory in Aging. PLoS ONE, 2012, 7, e45134. | 1.1 | 44 |
| 106 | Fine Mapping of Genetic Variants in BIN1, CLU, CR1 and PICALM for Association with Cerebrospinal Fluid Biomarkers for Alzheimer's Disease. PLoS ONE, 2011, 6, e15918. | 1.1 | 64 |
| 107 | Common variants at MS4A4/MS4A6E, CD2AP, CD33 and EPHA1 are associated with late-onset Alzheimer's disease. Nature Genetics, 2011, 43, 436-441. | 9.4 | 1,676 |
| 108 | Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. Nature Genetics, 2011, 43, 429-435. | 9.4 | 1,708 |

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|-----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 109 | Performances on the CogState and Standard Neuropsychological Batteries Among HIV Patients Without Dementia. <i>AIDS and Behavior</i> , 2011, 15, 1902-1909. | 1.4 | 52 |
| 110 | Association and Expression Analyses With Single-Nucleotide Polymorphisms in <i>TOMM40</i> in Alzheimer Disease. <i>Archives of Neurology</i> , 2011, 68, 1013. | 4.9 | 97 |
| 111 | SNPs Associated with Cerebrospinal Fluid Phospho-Tau Levels Influence Rate of Decline in Alzheimer's Disease. <i>PLoS Genetics</i> , 2010, 6, e1001101. | 1.5 | 111 |
| 112 | Genetic Evidence Implicates the Immune System and Cholesterol Metabolism in the Aetiology of Alzheimer's Disease. <i>PLoS ONE</i> , 2010, 5, e13950. | 1.1 | 347 |
| 113 | Validating predicted biological effects of Alzheimer's disease associated SNPs using CSF biomarker levels. <i>Journal of Alzheimer's Disease</i> , 2010, 21, 833-42. | 1.2 | 43 |
| 114 | Rs5848 Variant Influences <i>GRN</i> mRNA Levels in Brain and Peripheral Mononuclear Cells in Patients with Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2009, 18, 603-612. | 1.2 | 59 |
| 115 | Alzheimer's disease risk variants show association with cerebrospinal fluid amyloid beta. <i>Neurogenetics</i> , 2009, 10, 13-17. | 0.7 | 80 |
| 116 | Genome-wide association study identifies variants at <i>CLU</i> and <i>PICALM</i> associated with Alzheimer's disease. <i>Nature Genetics</i> , 2009, 41, 1088-1093. | 9.4 | 2,697 |
| 117 | Molecular characterization of novel progranulin (<i>GRN</i>) mutations in frontotemporal dementia. <i>Human Mutation</i> , 2008, 29, 512-521. | 1.1 | 71 |
| 118 | Novel presenilin 1 variant (P117A) causing Alzheimer's disease in the fourth decade of life. <i>Neuroscience Letters</i> , 2008, 438, 257-259. | 1.0 | 14 |
| 119 | Variation in <i>MAPT</i> is associated with cerebrospinal fluid tau levels in the presence of amyloid-beta deposition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 8050-8054. | 3.3 | 84 |
| 120 | Association studies between common variants in prolyl isomerase Pin1 and the risk for late-onset Alzheimer's disease. <i>Neuroscience Letters</i> , 2007, 419, 15-17. | 1.0 | 25 |
| 121 | Association studies testing for risk for late-onset Alzheimer's disease with common variants in the β^2 -amyloid precursor protein (<i>APP</i>). <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007, 144B, 469-474. | 1.1 | 11 |
| 122 | Extreme cerebrospinal fluid amyloid β^2 levels identify family with late-onset Alzheimer's disease presenilin 1 mutation. <i>Annals of Neurology</i> , 2007, 61, 446-453. | 2.8 | 87 |
| 123 | Haplotype-based association analysis of the <i>MAPT</i> locus in late onset Alzheimer's disease. <i>BMC Genetics</i> , 2007, 8, 3. | 2.7 | 45 |
| 124 | Apolipoprotein E levels in cerebrospinal fluid and the effects of <i>ABCA1</i> polymorphisms. <i>Molecular Neurodegeneration</i> , 2007, 2, 7. | 4.4 | 68 |
| 125 | Identification and validation of novel CSF biomarkers for early stages of Alzheimer's disease. <i>Proteomics - Clinical Applications</i> , 2007, 1, 1373-1384. | 0.8 | 66 |
| 126 | A Scan of Chromosome 10 Identifies a Novel Locus Showing Strong Association with Late-Onset Alzheimer Disease. <i>American Journal of Human Genetics</i> , 2006, 78, 78-88. | 2.6 | 157 |

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|-----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 127 | Ubiquilin 1 polymorphisms are not associated with late-onset Alzheimer's disease. <i>Annals of Neurology</i> , 2006, 59, 21-26. | 2.8 | 37 |
| 128 | HDDD2 is a familial frontotemporal lobar degeneration with ubiquitin-positive, tau-negative inclusions caused by a missense mutation in the signal peptide of progranulin. <i>Annals of Neurology</i> , 2006, 60, 314-322. | 2.8 | 186 |
| 129 | DAPK1 variants are associated with Alzheimer's disease and allele-specific expression. <i>Human Molecular Genetics</i> , 2006, 15, 2560-2568. | 1.4 | 125 |
| 130 | Association studies between risk for late-onset Alzheimer's disease and variants in insulin degrading enzyme. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2005, 136B, 62-68. | 1.1 | 35 |
| 131 | Microsatellites versus single-nucleotide polymorphisms in linkage analysis for quantitative and qualitative measures. <i>BMC Genetics</i> , 2005, 6, S122. | 2.7 | 15 |
| 132 | An analysis of identical single-nucleotide polymorphisms genotyped by two different platforms. <i>BMC Genetics</i> , 2005, 6, S152. | 2.7 | 6 |
| 133 | The efficacy of short tandem repeat polymorphisms versus single-nucleotide polymorphisms for resolving population structure. <i>BMC Genetics</i> , 2005, 6, S84. | 2.7 | 7 |
| 134 | Association of late-onset Alzheimer's disease with genetic variation in multiple members of the GAPD gene family. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 15688-15693. | 3.3 | 134 |
| 135 | Phylogeographic and nested clade analysis of the stonefly <i>Pteronarcys californica</i> (Plecoptera:Pteronarcyidae) in the western USA. <i>Journal of the North American Benthological Society</i> , 2004, 23, 824-838. | 3.0 | 27 |
| 136 | Genome assembly of the roundjaw bonefish (<i>Albula glossodonta</i>), a vulnerable circumtropical sportfish. <i>GigaByte</i> , 0, 2022, 1-29. | 0.0 | 1 |